RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/587,371
Source: IFWP
Date Processed by STIC: 08/07/2006

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IFWP

RAW SEQUENCE LISTING DATE: 08/07/2006
PATENT APPLICATION: **US/10/587,371** TIME: 10:04:27

Input Set : **F:\AMBX-002800US.ST25.txt**
Output Set: **N:\CRF4\08072006\J587371.raw**

3 <110> APPLICANT: Cho, Ho S
4 Daniel, Thomas
5 DiMarchi, Richard
6 Hays, Anna-Maria
7 Wilson, Troy
8 Sim, Bee-Cheng
9 Litzinger, David
11 <120> TITLE OF INVENTION: Modified Human Four Helical Bundle Polypeptides and Their Uses
13 <130> FILE REFERENCE: AMBX-0028.00PCT
C--> 15 <140> CURRENT APPLICATION NUMBER: US/10/587,371
C--> 15 <141> CURRENT FILING DATE: 2006-07-26
15 <150> PRIOR APPLICATION NUMBER: 60/541,528
16 <151> PRIOR FILING DATE: 2004-02-02
18 <150> PRIOR APPLICATION NUMBER: 60/581,314
19 <151> PRIOR FILING DATE: 2004-06-18
21 <150> PRIOR APPLICATION NUMBER: 60/581,175
22 <151> PRIOR FILING DATE: 2004-06-18
24 <150> PRIOR APPLICATION NUMBER: 60/580,885
25 <151> PRIOR FILING DATE: 2004-06-18
27 <150> PRIOR APPLICATION NUMBER: 60/638,616
28 <151> PRIOR FILING DATE: 2004-12-22
30 <160> NUMBER OF SEQ ID NOS: 46
32 <170> SOFTWARE: PatentIn version 3.3
34 <210> SEQ ID NO: 1
35 <211> LENGTH: 217
36 <212> TYPE: PRT
37 <213> ORGANISM: Homo sapiens
39 <400> SEQUENCE: 1

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45 Cys Leu Pro Trp Leu Gln Glu Gly Ser Ala Phe Pro Thr Ile Pro Leu
46             20                  25                  30
49 Ser Arg Leu Phe Asp Asn Ala Met Leu Arg Ala His Arg Leu His Gln
50         35                  40                  45
53 Leu Ala Phe Asp Thr Tyr Gln Glu Phe Glu Glu Ala Tyr Ile Pro Lys
54     50                  55                  60
57 Glu Gln Lys Tyr Ser Phe Leu Gln Asn Pro Gln Thr Ser Leu Cys Phe
58 65                  70                  75                  80
61 Ser Glu Ser Ile Pro Thr Pro Ser Asn Arg Glu Glu Thr Gln Gln Lys
62                 85                  90                  95
65 Ser Asn Leu Glu Leu Leu Arg Ile Ser Leu Leu Leu Ile Gln Ser Trp
66             100                 105                 110
69 Leu Glu Pro Val Gln Phe Leu Arg Ser Val Phe Ala Asn Ser Leu Val
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70          115                 120                 125
73 Tyr Gly Ala Ser Asp Ser Asn Val Tyr Asp Leu Leu Lys Asp Leu Glu
74      130                 135                 140
77 Glu Gly Ile Gln Thr Leu Met Gly Arg Leu Glu Asp Gly Ser Pro Arg
78 145                 150                 155                 160
81 Thr Gly Gln Ile Phe Lys Gln Thr Tyr Ser Lys Phe Asp Thr Asn Ser
82                  165                 170                 175
85 His Asn Asp Asp Ala Leu Leu Lys Asn Tyr Gly Leu Leu Tyr Cys Phe
86              180                 185                 190
89 Arg Lys Asp Met Asp Lys Val Glu Thr Phe Leu Arg Ile Val Gln Cys
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99 <212> TYPE: PRT
100 <213> ORGANISM: Homo sapiens
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108 Ala His Arg Leu His Gln Leu Ala Phe Asp Thr Tyr Gln Glu Phe Glu
109             20                  25                  30
112 Glu Ala Tyr Ile Pro Lys Glu Gln Lys Tyr Ser Phe Leu Gln Asn Pro
113         35                  40                  45
116 Gln Thr Ser Leu Cys Phe Ser Glu Ser Ile Pro Thr Pro Ser Asn Arg
117     50                  55                  60
120 Glu Glu Thr Gln Gln Lys Ser Asn Leu Glu Leu Leu Arg Ile Ser Leu
121 65                  70                  75                  80
124 Leu Leu Ile Gln Ser Trp Leu Glu Pro Val Gln Phe Leu Arg Ser Val
125                 85                  90                  95
128 Phe Ala Asn Ser Leu Val Tyr Gly Ala Ser Asp Ser Asn Val Tyr Asp
129             100                 105                 110
132 Leu Leu Lys Asp Leu Glu Glu Gly Ile Gln Thr Leu Met Gly Arg Leu
133         115                 120                 125
136 Glu Asp Gly Ser Pro Arg Thr Gly Gln Ile Phe Lys Gln Thr Tyr Ser
137     130                 135                 140
140 Lys Phe Asp Thr Asn Ser His Asn Asp Asp Ala Leu Leu Lys Asn Tyr
141 145                 150                 155                 160
144 Gly Leu Leu Tyr Cys Phe Arg Lys Asp Met Asp Lys Val Glu Thr Phe
145                 165                 170                 175
148 Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
149             180                 185                 190
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154 <212> TYPE: PRT
155 <213> ORGANISM: Homo sapiens
157 <400> SEQUENCE: 3
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160 1               5                   10                  15
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```
163 Ala His Arg Leu His Gln Leu Ala Phe Asp Thr Tyr Gln Glu Phe Asn
164             20                  25                  30
167 Pro Gln Thr Ser Leu Cys Phe Ser Glu Ser Ile Pro Thr Pro Ser Asn
168         35                  40                  45
171 Arg Glu Glu Thr Gln Gln Lys Ser Asn Leu Glu Leu Leu Arg Ile Ser
172     50                  55                  60
175 Leu Leu Leu Ile Gln Ser Trp Leu Glu Pro Val Gln Phe Leu Arg Ser
176 65                  70                  75                  80
179 Val Phe Ala Asn Ser Leu Val Tyr Gly Ala Ser Asp Ser Asn Val Tyr
180                 85                  90                  95
183 Asp Leu Leu Lys Asp Leu Glu Glu Gly Ile Gln Thr Leu Met Gly Arg
184             100                 105                 110
187 Leu Glu Asp Gly Ser Pro Arg Thr Gly Gln Ile Phe Lys Gln Thr Tyr
188         115                 120                 125
191 Ser Lys Phe Asp Thr Asn Ser His Asn Asp Asp Ala Leu Leu Lys Asn
192     130                 135                 140
195 Tyr Gly Leu Leu Tyr Cys Phe Arg Lys Asp Met Asp Lys Val Glu Thr
196 145                 150                 155                 160
199 Phe Leu Arg Ile Val Gln Cys Arg Ser Val Glu Gly Ser Cys Gly Phe
200                 165                 170                 175
203 <210> SEQ ID NO: 4
204 <211> LENGTH: 77
205 <212> TYPE: DNA
206 <213> ORGANISM: Methanococcus jannaschii
208 <400> SEQUENCE: 4
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215 <211> LENGTH: 88
216 <212> TYPE: DNA
217 <213> ORGANISM: Halobacterium sp. NRC-1
219 <400> SEQUENCE: 5
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226 <211> LENGTH: 89
227 <212> TYPE: DNA
228 <213> ORGANISM: Halobacterium sp. NRC-1
230 <400> SEQUENCE: 6
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237 <211> LENGTH: 306
238 <212> TYPE: PRT
239 <213> ORGANISM: Methanococcus jannaschii
241 <400> SEQUENCE: 7
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```
248                 20                  25                  30
251 Ile Gly Phe Glu Pro Ser Gly Lys Ile His Leu Gly His Tyr Leu Gln
252         35                  40                  45
255 Ile Lys Lys Met Ile Asp Leu Gln Asn Ala Gly Phe Asp Ile Ile Ile
256     50                  55                  60
259 Leu Leu Ala Asp Leu His Ala Tyr Leu Asn Gln Lys Gly Glu Leu Asp
260 65                  70                  75                  80
263 Glu Ile Arg Lys Ile Gly Asp Tyr Asn Lys Lys Val Phe Glu Ala Met
264                 85                  90                  95
267 Gly Leu Lys Ala Lys Tyr Val Tyr Gly Ser Thr Phe Gln Leu Asp Lys
268             100                 105                 110
271 Asp Tyr Thr Leu Asn Val Tyr Arg Leu Ala Leu Lys Thr Thr Leu Lys
272         115                 120                 125
275 Arg Ala Arg Arg Ser Met Glu Leu Ile Ala Arg Glu Asp Glu Asn Pro
276     130                 135                 140
279 Lys Val Ala Glu Val Ile Tyr Pro Ile Met Gln Val Asn Thr Tyr Tyr
280 145                 150                 155                 160
283 Tyr Leu Gly Val Asp Val Ala Val Gly Gly Met Glu Gln Arg Lys Ile
284                 165                 170                 175
287 His Met Leu Ala Arg Glu Leu Leu Pro Lys Lys Val Val Cys Ile His
288             180                 185                 190
291 Asn Pro Val Leu Thr Gly Leu Asp Gly Glu Gly Lys Met Ser Ser Ser
292         195                 200                 205
295 Lys Gly Asn Phe Ile Ala Val Asp Asp Ser Pro Glu Glu Ile Arg Ala
296     210                 215                 220
299 Lys Ile Lys Lys Ala Tyr Cys Pro Ala Gly Val Val Glu Gly Asn Pro
300 225                 230                 235                 240
303 Ile Met Glu Ile Ala Lys Tyr Phe Leu Glu Tyr Pro Leu Thr Ile Lys
304                 245                 250                 255
307 Arg Pro Glu Lys Phe Gly Gly Asp Leu Thr Val Asn Ser Tyr Glu Glu
308             260                 265                 270
311 Leu Glu Ser Leu Phe Lys Asn Lys Glu Leu His Pro Met Asp Leu Lys
312         275                 280                 285
315 Asn Ala Val Ala Glu Glu Leu Ile Lys Ile Leu Glu Pro Ile Arg Lys
316     290                 295                 300
319 Arg Leu
320 305
323 <210> SEQ ID NO: 8
324 <211> LENGTH: 306
325 <212> TYPE: PRT
326 <213> ORGANISM: Methanococcus jannaschii
328 <400> SEQUENCE: 8
330 Met Asp Glu Phe Glu Met Ile Lys Arg Asn Thr Ser Glu Ile Ile Ser
331 1               5                   10                  15
334 Glu Glu Glu Leu Arg Glu Val Leu Lys Lys Asp Glu Lys Ser Ala Gly
335             20                  25                  30
338 Ile Gly Phe Glu Pro Ser Gly Lys Ile His Leu Gly His Tyr Leu Gln
339         35                  40                  45
342 Ile Lys Lys Met Ile Asp Leu Gln Asn Ala Gly Phe Asp Ile Ile Ile
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```
343     50                  55                  60
346 Leu Leu Ala Asp Leu His Ala Tyr Leu Asn Gln Lys Gly Glu Leu Asp
347 65                  70                  75                  80
350 Glu Ile Arg Lys Ile Gly Asp Tyr Asn Lys Lys Val Phe Glu Ala Met
351                 85                  90                  95
354 Gly Leu Lys Ala Lys Tyr Val Tyr Gly Ser Ser Phe Gln Leu Asp Lys
355             100                 105                 110
358 Asp Tyr Thr Leu Asn Val Tyr Arg Leu Ala Leu Lys Thr Thr Leu Lys
359         115                 120                 125
362 Arg Ala Arg Arg Ser Met Glu Leu Ile Ala Arg Glu Asp Glu Asn Pro
363     130                 135                 140
366 Lys Val Ala Glu Val Ile Tyr Pro Ile Met Gln Val Asn Thr Ser His
367 145                 150                 155                 160
370 Tyr Leu Gly Val Asp Val Ala Val Gly Gly Met Glu Gln Arg Lys Ile
371                 165                 170                 175
374 His Met Leu Ala Arg Glu Leu Leu Pro Lys Lys Val Val Cys Ile His
375             180                 185                 190
378 Asn Pro Val Leu Thr Gly Leu Asp Gly Glu Gly Lys Met Ser Ser Ser
379         195                 200                 205
382 Lys Gly Asn Phe Ile Ala Val Asp Asp Ser Pro Glu Glu Ile Arg Ala
383     210                 215                 220
386 Lys Ile Lys Lys Ala Tyr Cys Pro Ala Gly Val Val Glu Gly Asn Pro
387 225                 230                 235                 240
390 Ile Met Glu Ile Ala Lys Tyr Phe Leu Glu Tyr Pro Leu Thr Ile Lys
391                 245                 250                 255
394 Arg Pro Glu Lys Phe Gly Gly Asp Leu Thr Val Asn Ser Tyr Glu Glu
395             260                 265                 270
398 Leu Glu Ser Leu Phe Lys Asn Lys Glu Leu His Pro Met Asp Leu Lys
399         275                 280                 285
402 Asn Ala Val Ala Glu Glu Leu Ile Lys Ile Leu Glu Pro Ile Arg Lys
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406 Arg Leu
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410 <210> SEQ ID NO: 9
411 <211> LENGTH: 305
412 <212> TYPE: PRT
413 <213> ORGANISM: Methanococcus jannaschii
415 <400> SEQUENCE: 9
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422             20                  25                  30
425 Ile Gly Phe Glu Pro Ser Gly Lys Ile His Leu Gly His Tyr Leu Gln
426         35                  40                  45
429 Ile Lys Lys Met Ile Asp Leu Gln Asn Ala Gly Phe Asp Ile Ile Ile
430     50                  55                  60
433 Leu Leu Ala Asp Leu His Ala Tyr Leu Asn Gln Lys Gly Glu Leu Asp
434 65                  70                  75                  80
437 Glu Ile Arg Lys Ile Gly Asp Tyr Asn Lys Lys Val Phe Glu Ala Met
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VERIFICATION SUMMARY DATE: 08/07/2006
PATENT APPLICATION: **US/10/587,371** TIME: 10:04:28

Input Set : **F:\AMBX-002800US.ST25.txt**
Output Set: **N:\CRF4\08072006\J587371.raw**

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L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date